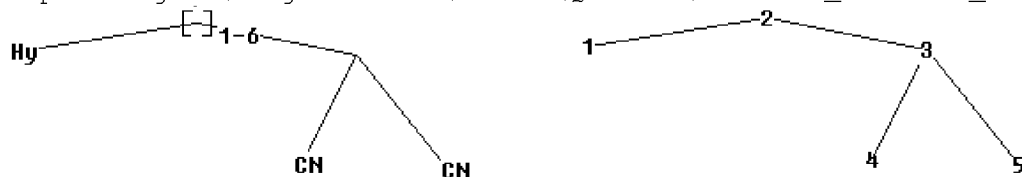


=> FILE REGISTRY

=>

Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_2.str



chain nodes :

1 2 3 4 5

chain bonds :

1-2 2-3 3-4 3-5

exact/norm bonds :

1-2

exact bonds :

2-3 3-4 3-5

Match level :

1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS

Generic attributes :

1:

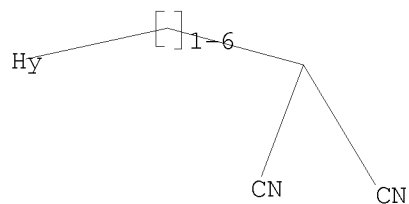
Number of Hetero Atoms : 2 or more

L1 STRUCTURE UPLOADED

=> d

L1 HAS NO ANSWERS

L1 STR



Structure attributes must be viewed using STN Express query preparation.

=> s l1 full

FULL SEARCH INITIATED 06:46:21 FILE 'REGISTRY'

FULL SCREEN SEARCH COMPLETED - 19112 TO ITERATE

100.0% PROCESSED 19112 ITERATIONS

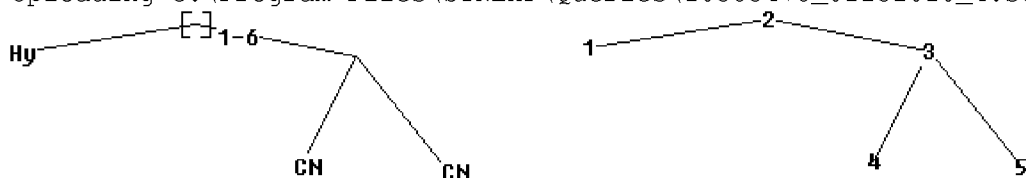
1000 ANSWERS

SEARCH TIME: 00.00.01

L3 1000 SEA SSS FUL L1

=>

Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_4.str



chain nodes :

1 2 3 4 5

chain bonds :

1-2 2-3 3-4 3-5

exact/norm bonds :

1-2

exact bonds :

2-3 3-4 3-5

Match level :

1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS

Generic attributes :

1:

Saturation : Unsaturated

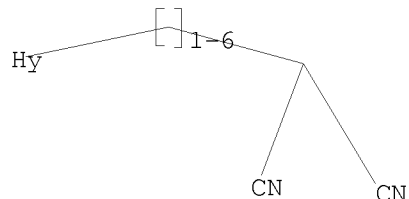
Number of Hetero Atoms : 2 or more

L8 STRUCTURE UPLOADED

=> d

L8 HAS NO ANSWERS

L8 STR



Structure attributes must be viewed using STN Express query preparation.

=> s 18 sss sub=16 full

FULL SUBSET SEARCH INITIATED 06:49:34 FILE 'REGISTRY'

FULL SUBSET SCREEN SEARCH COMPLETED - 520 TO ITERATE

100.0% PROCESSED 520 ITERATIONS

488 ANSWERS

SEARCH TIME: 00.00.01

L10 488 SEA SUB=L6 SSS FUL L8

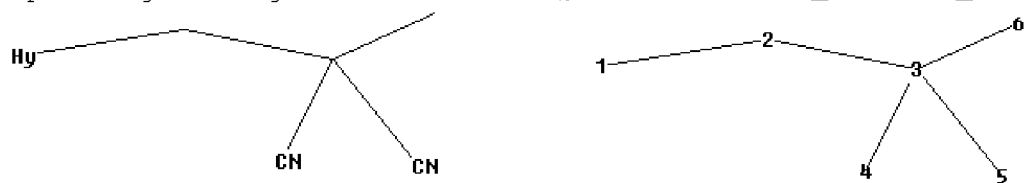
=> s 110 and caplus/lc

69979096 CAPLUS/LC

L11 287 L10 AND CAPLUS/LC

=>

Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_5.str



chain nodes :

1 2 3 4 5

ring/chain nodes :

6

chain bonds :

1-2 2-3 3-4 3-5 3-6

exact/norm bonds :

1-2

exact bonds :

2-3 3-4 3-5 3-6

Match level :

1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS

Generic attributes :

1:

Saturation : Unsaturated

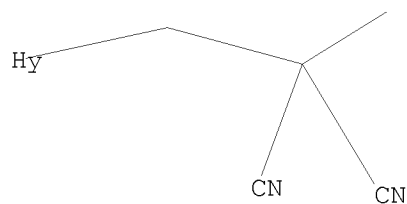
Number of Hetero Atoms : 2 or more

L13 STRUCTURE UPLOADED

=> d

L13 HAS NO ANSWERS

L13 STR



=> s l13 sss sub=l10 full

FULL SUBSET SEARCH INITIATED 06:50:47 FILE 'REGISTRY'

FULL SUBSET SCREEN SEARCH COMPLETED - 153 TO ITERATE

100.0% PROCESSED 153 ITERATIONS

132 ANSWERS

SEARCH TIME: 00.00.01

L15 132 SEA SUB=L10 SSS FUL L13

=> s l15 and caplus/lc

69979096 CAPLUS/LC
L16 132 L15 AND CAPLUS/LC

=> file caplus

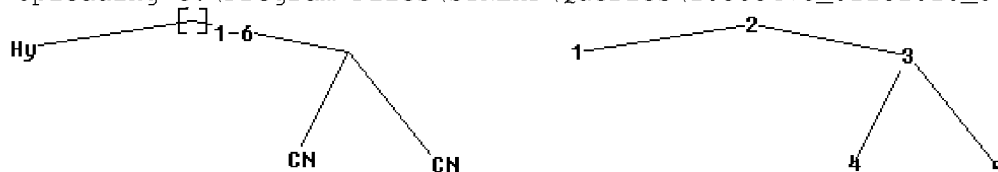
=> s l16

L17 10 L16

=> d l17 ibib gi abs hitstr 1-10

=>

Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_6.str



chain nodes :

1 2 3 4 5

chain bonds :

1-2 2-3 3-4 3-5

exact/norm bonds :

1-2

exact bonds :

2-3 3-4 3-5

Match level :

1:Atom 2:CLASS 3:CLASS 4:CLASS 5:CLASS

Generic attributes :

1:

Saturation : Unsaturated

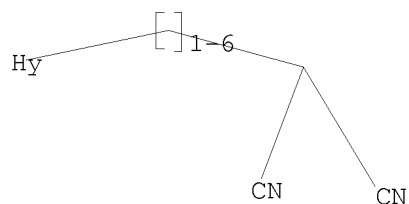
Number of Hetero Atoms : 2 or more

L19 STRUCTURE UPLOADED

=> d

L19 HAS NO ANSWERS

L19 STR



=> s l19 sss sub=l18 full

FULL SUBSET SEARCH INITIATED 07:00:43 FILE 'REGISTRY'

FULL SUBSET SCREEN SEARCH COMPLETED - 742 TO ITERATE

100.0% PROCESSED 742 ITERATIONS

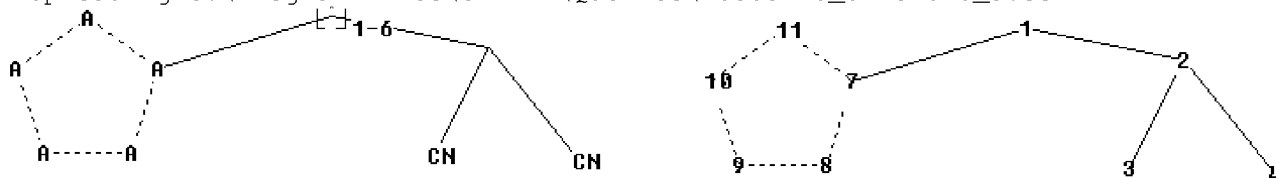
690 ANSWERS

SEARCH TIME: 00.00.01

L21 690 SEA SUB=L18 SSS FUL L19

=>

Uploading C:\Program Files\STNEXP\Queries\10585478_01252010_3.str



chain nodes :

1 2 3 4

ring nodes :

7 8 9 10 11

chain bonds :

1-2 1-7 2-3 2-4

ring bonds :

7-8 7-11 8-9 9-10 10-11

exact/norm bonds :

1-7 7-8 7-11 8-9 9-10 10-11

exact bonds :

1-2 2-3 2-4

Match level :

1:CLASS 2:CLASS 3:CLASS 4:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom

L22 STRUCTURE UPLOADED

=> s l22 sss sub=l21 full

FULL SUBSET SEARCH INITIATED 07:01:31 FILE 'REGISTRY'

FULL SUBSET SCREEN SEARCH COMPLETED - 690 TO ITERATE

100.0% PROCESSED 690 ITERATIONS

287 ANSWERS

SEARCH TIME: 00.00.01

L24 287 SEA SUB=L21 SSS FUL L22

=> file caplus

=> s l24/USES

83 L24

7936248 USES/RL

L25 13 L24/USES

(L24 (L) USES/RL)

=> d l25 ibib gi abs hitstr 1-13

=> logoff hold